



OIPE

RAW SEQUENCE LISTING DATE: 02/07/2002 PATENT APPLICATION: US/09/888,243 TIME: 13:33:08

Input Set : A:\01997.211003.SEQLIST.TXT Output Set: N:\CRF3\02072002\1888243.raw

```
4 <110> APPLICANT: Horvitz, H. Robert
 5
         Yuan, Junying
 6
         Shaham, Shai
   <120> TITLE OF INVENTION: Relatedness of Human Interleukin-1beta
         Convertase Gene to a C. Elegans Cell Death Gene, Inhibitory
10
         Portions of these Genes and Uses Therefor
13 <130> FILE REFERENCE: 01997/211003
15 <140> CURRENT APPLICATION NUMBER: US 09/888,243
16 <141> CURRENT FILING DATE: 2001-06-22
18 <150> PRIOR APPLICATION NUMBER: US 09/083,662
19 <151> PRIOR FILING DATE: 1998-05-22
21 <150> PRIOR APPLICATION NUMBER: US 08/394,189
22 <151> PRIOR FILING DATE: 1995-02-24
24 <150> PRIOR APPLICATION NUMBER: US 08/282,211
                                                        ENTERED
25 <151> PRIOR FILING DATE: 1994-07-11
27 <150> PRIOR APPLICATION NUMBER: US 07/984,182
28 <151> PRIOR FILING DATE: 1992-11-20
30 <150> PRIOR APPLICATION NUMBER: US 07/897,788
31 <151> PRIOR FILING DATE: 1992-06-12
33 <160> NUMBER OF SEQ ID NOS: 30
35 <170> SOFTWARE: FastSEQ for Windows Version 4.0
37 <210> SEQ ID NO: 1
38 <211> LENGTH: 7653
39 <212> TYPE: DNA
40 <213> ORGANISM: Caenorhabditis elegans
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45 ttgtcgaatt aatatcccta ttatcacttt ttcatgctca tcctcgagcg gcacgtcctc 180
46 aaagaattgt gagagcaaac gegeteeeat tgaceteeac acteageege caaaacaaac 240
47 gttcgaacat tcgtgtgttg tgctcctttt ccgttatctt gcagtcatct tttgtcgttt 300
48 ttttctttgt tctttttgtt gaacgtgttg ctaagcaatt attacatcaa ttgaagaaaa 360
49 ggctcgccga tttattgttg ccagaaagat tctgagattc tcgaaqtcga ttttataata 420
50 tttaacettg gtttttgcat tgtttegttt aaaaaaaeca etgtttatgt gaaaaaegat 480
51 tagtttacta ataaaactac ttttaaacct ttacctttac ctcaccgctc cgtgttcatg 540
52 gctcatagat tttcgatact caaatccaaa aataaattta cgagggcaat taatgtgaaa 600
53 caaaaacaat cctaagattt ccacatgttt gacctctccg gcaccttctt ccttagcccc 660
54 accactccat cacctctttg gcggtgttct tcgaaaccca cttaggaaag cagtgtgtat 720
55 ctcatttggt atgetetttt egattttata getetttgte geaattteaa tgetttaaac 780
56 aatccaaatc gcattatatt tgtgcatgga ggcaaatgac ggggttggaa tcttagatga 840
57 gatcaggagc tttcagggta aacgcccggt tcattttgta ccacatttca tcattttcct 900
58 gtcgtccttg gtatcctcaa cttgtcccgg ttttgttttc ggtacactct tccgtgatgc 960
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59 cacctgtctc cgtctcaatt atcgtttaga aatgtgaact gtccagatgg gtgactcata 1020

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60 ttgctgctgc tacaatccac tttcttttct catcggcagt cttacgagcc catcataaac 1080 61 tttttttcc gcgaaatttg caataaaccg gccaaaaact ttctccaaat tgttacgcaa 1140 62 tatatacaat ccataagaat atcttctcaa tgtttatgat ttcttcgcag cactttctct 1200 63 togtqtqcta acatottatt tttataatat ttooqotaaa attooqattt ttgagtatta 1260 64 atttatcgta aaattatcat aatagcaccg aaaactacta aaaatggtaa aagctccttt 1320 65 taaatcggct cgacattatc gtattaagga atcacaaaat tctgagaatg cgtactgcgc 1380 66 aacatatttg acggcaaaat atctcgtagc gaaaactaca gtaattcttt aaatgactac 1440 67 tgtagcgctt gtgtcgattt acgggctcaa tttttgaaaa taatttttt tttcgaattt 1500 68 tgataacccg taaatcgtca caacgctaca gtagtcattt aaaggattac tgtagttcta 1560 69 gctacgagat attttgcgcg ccaaatatga ctgtaatacg cattctctga attttgtgtt 1620 70 teegtaataa titeacaaga tittggeatt eeactitaaa ggegeacagg attiatteea 1680 71 atgggtctcg gcacgcaaaa agtttgatag acttttaaat tctccttgca tttttaattc 1740 72 aattactaaa attttcgtga atttttctgt taaaattttt aaaatcagtt ttctaatatt 1800 73 ttccaggctg acaaacagaa acaaaaacac aacaaacatt ttaaaaaatca gttttcaaat 1860 74 taaaaataac gatttctcat tgaaaattgt gttttatgtt tgcgaaaata aaagagaact 1920 75 gattcaaaac aattttaaca aaaaaaaacc ccaaaattcg ccagaaatca agataaaaaa 1980 76 ttcaagaggg tcaaaatttt ccqattttac tgactttcac cttttttttc gtagttcagt 2040 77 gcagttgttg gagtttttga cgaaaactag gaaaaaaatc gataaaaatt actcaaatcg 2100 78 agctgaattt tgaggacaat gtttaaaaaa aaacactatt tttccaataa tttcactcat 2160 79 tttcagacta aatcgaaaat caaatcgtac tctgactacg ggtcagtaga gaggtcaacc 2220 80 atcagcogaa gatgatgogt caagatagaa ggagottgot agagaggaac attatgatgt 2280 81 tetetagtea tetaaaagte gatgaaatty tegaagttet eategeaaaa caagtgttga 2340 82 atagtgataa tggagatatg attaatgtga gtttttaatc gaataataat tttaaaaaaa 2400 83 aattgataat ataaagaata tttttgcagt catgtggaac ggttcgcgag aagagacggg 2460 84 agatcgtgaa agcagtgcaa cgacggrgag atgtggcgtt cgacgcgttt tatgatgctc 2520 85 ttcgctctac gggacacgaa ggacttgctg aagttcttga acctctcgcc agatcgtagg 2580 86 tttttaaaqt toqqoqoaaa aqoaaqqqto toacqqaaaa aagaggogga toqtaatttt 2640 87 gcaacccacc ggcacggttt tttcctccga aaatcggaaa ttatgcactt tcccaaatat 2700 88 ttgaagtgaa atatatttta tttactgaaa gctcgagtga ttatttattt tttaacacta 2760 89 attttcgtgg cgcaaaaggc cattttgtag atttgccgaa aatacttgtc acacacac 2820 90 acacacatct ccttcaaata tccctttttc cagtgttgac tcgaatgctg tcgaattcga 2880 91 gtgtccaatg tcaccggcaa gccatcgtcg gagccgcgca ttgagccccg ccggctacac 2940 92 ttcaccgacc cgagttcacc gtgacagcgt ctcttcagtg tcatcattca cttcttatca 3000 93 ggatatetae teaagageaa gatetegtte tegategegt geaetteatt categgateg 3060 94 acacaattat tcatctcctc cagtcaacge atttcccage caacettgta tgttgatgcg 3120 95 aacactaaat totgagaatg ogcattacto aacatatttg acgogoaaat atotogtago 3180 96 gaaaaataca gtaaccettt aaatgactat tgtagtgtcg atttacgggc tcgattttcg 3240 97 aaacgaatat atgctcgaat tgtgacaacg aattttaatt tgtcattttt gtgttttctt 3300 98 ttgatatttt tgatcaatta ataaattatt teegtaaaca gacaceageg etaeagtaet 3360 99 cttttaaaga gttacagtag ttttcgcttc aagatatttt gaaaagaatt ttaaacattt 3420 100 tgaaaaaaa tcatctaaca tgtgccaaaa cgcttttttc aagtttcgca gattttttga 3480 101 tttttttcat tcaagatatg cttattaaca catataatta tcattaatgt gaatttcttg 3540 102 tagaaatttt gggcttttcg ttctagtatg ctctactttt gaaattgctc aacgaaaaaa 3600 103 tcatqtgqtt tgttcatatg aatgacgaaa aatagcaatt ttttatatat tttcccctat 3660 104 tcatgttgtg cagaaaaata gtaaaaaagc gcatgcattt ttcgacattt tttacatcga 3720 105 acgacagete actteacatg etgaagaega gagaegegga gaaataecae acatetttet 3780 106 gcgtctctcg tcttcagcat gtgaaatggg atctcggtcg atgtaaaaaa atgtcgaata 3840 107 atgtaaaaaa tgcatgcgtt tttttacact tttctgcaca aatgaatagg gggaaaatgt 3900 108 attaaaatac attttttgta tttttcaaca tcacatgatt aaccccatta ttttttcgtt 3960

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110 tttattgata attatagatg ttaataagca tatcttgaat gaaagtcagc aaaaatatgt 4080
111 gcgaaacacc tgaaaaaaat caaaaattct gcgaaaattg aaaaaatgca ttaaaataca 4140
112 tttttgcatt tttctacatc acatgaatgt agaaaattaa aagggaaatc aaaatttcta 4200
113 gaggatataa ttgaatgaaa cattgcgaaa ttaaaatgtg cgaaacgtca aaaaagagga 4260
114 aatttgggta tcaaaatcga tcctaaaacc aacacatttc agcatccgcc aactcttcat 4320
115 tcaccqqatq ctcttctctc qqatacaqtt caaqtcqtaa tcgctcattc agcaaagctt 4380
116 ctggaccaac tcaatacata ttccatgaag aggatatgaa ctttgtcgat gcaccaacca 4440
117 taaqccqtqt tttcqacqaq aaaaccatqt acaqaaactt ctcqaqtcct cqtqqaatqt 4500
118 gcctcatcat aaataatgaa cactttgagc agatgccaac acggaatggt accaaggccg 4560
119 acaaqqacaa tottaccaat ttqttcagat gcatqqqcta tacqqttatt tqcaaqqaca 4620
120 atctgacggg aagggtacgg cgaaattata ttacccaaac gcgaaatttg ccattttgcg 4680
121 ccgaaaatgt ggcgcccggt ctcgacacga caatttgtgt taaatgcaaa aatgtataat 4740
122 tttgcaaaaa acaaaatttt gaacttccgc gaaaatgatt tacctagttt cgaaattttc 4800
123 gttttttccg gctacattat gtgtttttc ttagtttttc tataatattt gatgtaaaaa 4860
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125 ttctqaattt tcaaaattat ccaaaaatqc acaatttaaa atttqtqaaa attggcaaac 4980
126 ggtgtttcaa tatgaaatgt atttttaaaa actttaaaaa ccactccgga aaagcaataa 5040
127 aaatcaaaac aacgtcacaa ttcaaattca aaagttattc atccgatttg tttatttttg 5100
128 caaaatttga aaaaatcatg aaggatttag aaaagtttta taacattttt totagatttt 5160
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132 tggacaataa accttcctaa tcaccaaaaa gtaaaattga aatcttcgaa aagccaaaaa 5400
133 attcaaaaaa aaagtcgaat ttcgattttt tttttggttt tttggtccca aaaaccaaaa 5460
134 aaatcaattt totgoaaaat accaaaaaga aaccogaaaa aatttoocag cottgttoot 5520
135 aatgtaaact gatatttaat ttccagggaa tgctcctgac aattcgagac tttgccaaac 5580
136 acquatcaca cggagattct gcgatactcg tgattctatc acacggagaa gagaatgtga 5640
137 ttattggagt tgatgatata ccgattagta cacacgagat atatgatctt ctcaacgcgg 5700
138 caaatgctcc ccgtctggcg aataagccga aaatcgtttt tgtgcaggct tgtcgargcg 5760
139 qttcqttttt tattttaatt ttaatataaa tattttaaat aaattcattt tcagaacgtc 5820
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141 gatgggacaa tcgagacggg ccattgttca attttcttgg atgtgtgcgg ccgcaagtty 5940
142 aggttgcaat ttaatttctt gaatgagaat attccttcaa aaaatctaaa atagattttt 6000
143 attocagaaa gtoccgatog aaaaattgog atataattac gaaatttgtg ataaaatgac 6060
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151 gtcgyttgtg gatttcagac atcacaggga tcgaatattt tgaaacagat gccaraggta 6540
152 cttgaaacaa acaatgcatg tctaactttt aaggacacag aaaaataggc agaggctcct 6600
153 tttgcaagcc tgccgcgcgt caacctagaa ttttagtttt tagctaaaat gattgatttt 6660
154 qaatatttta tgctaatttt tttgcgttaa attttgaaat agtcactatt tatcgggttt 6720
155 ccaqtaaaaa atgtttatta gccattggat tttactgaaa acgaaaattt gtagtttttc 6780
156 aacgaaattt atcgattttt aaatgtaaaa aaaaatagcg aaaattacat caaccatcaa 6840
157 gcatttaagc caaaattgtt aactcattta aaaattaatt caaagttgtc cacgagtatt 6900
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Input Set: A:\01997.211003.SEQLIST.TXT
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    159 gaaaacaagg gatcggttta gatttttccc caaaatttaa attaaatttc agatgacaty 7020
    160 ccgcctgctc aaaaagttct acttttggcc ggaagcacga aactctgccg tctaaaattc 7080
    161 actogtgatt cattgoccaa ttgataattg totgtatott otoccccagt tototttogo 7140
    162 ccaattagtt taaaaccatg tgtatattgt tatcctatac tcatttcact ttatcattct 7200
    163 atcatttctc ttcccatttt cacacatttc catttctcta cgataatcta aaattatgac 7260
    164 gtttgtgtct cgaacgcata ataattttaa taactcgttt tgaatttgat tagttgttgt 7320
    165 gcccagtata tatgtatgta ctatgcttct atcaacaaaa tagtttcata gatcatcacc 7380
    166 ccaaccccac caacctaccg taccatattc atttttgccg ggaatcaatt tcgattaatt 7440
    167 ttaacctatt ttttcgccac aaaaaatcta atatttgaat taacgaatag cattcccatc 7500
    168 tctcccgtgc cggaatgcct cccggccttt taaagttcgg aacatttggc aattatgtat 7560
    169 aaatttgtag gtcccccca tcatttcccg cccatcatct caaattgcat tctttttcg 7620
    170 ccgtgatatc ccgattctgg tcagcaaaga tct
    172 <210> SEQ ID NO: 2
    173 <211> LENGTH: 503
    174 <212> TYPE: PRT
    175 <213> ORGANISM: Caenorhabditis elegans
    177 <220> FEATURE:
                                     ( , (
    178 <221> NAME/KEY: VARIANT /
    179 <222> LOCATION: 27, 65, 360, 403, 412, 428, 449, 466, 483, 486
    180 <223> OTHER INFORMATION: Xaa = Any Amino Acid
    182 <400> SEQUENCE: 2
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    184 1
W--> 185 Phe Ser Ser His Leu Lys Val Asp Glu Ile Xaa Glu Val Leu Ile Ala
                   20
                                        25
    187 Lys Gln Val Leu Asn Ser Asp Asn Gly Asp Met Ile Asn Ser Cys Gly
                                    40
               35
    189 Thr Val Arg Glu Lys Arg Arg Glu Ile Val Lys Ala Val Gln Arg Arg
    190 \sim 50
                                55
W--> 191 Xaa Asp Val Ala Phe Asp Ala Phe Tyr Asp Ala Leu Arg Ser Thr Gly
                                                75
    192 65
                            70
     193 His Glu Gly Leu Ala Glu Val Leu Glu Pro Leu Ala Arg Ser Val Asp
                                            90
     195 Ser Asn Ala Val Glu Phe Glu Cys Pro Met Ser Pro Ala Ser His Arg
                                        105
                    100
    197 Arg Ser Arg Ala Leu Ser Pro Ala Gly Tyr Thr Ser Pro Thr Arg Val
               115
                                    120
    199 His Arg Asp Ser Val Ser Ser Val Ser Ser Phe Thr Ser Tyr Gln Asp
                                135
                                                    140
    201 Ile Tyr Ser Arg Ala Arg Ser Arg Ser Arg Ser Arg Ala Leu His Ser
                                               155
                            150
    203 Ser Asp Arg His Asn Tyr Ser Ser Pro Pro Val Asn Ala Phe Pro Ser
                                            170
                        165
    205 Gln Pro Ser Ser Ala Asn Ser Ser Phe Thr Gly Cys Ser Ser Leu Gly
                                        185
                                                            190
                   180
    206
     207 Tyr Ser Ser Ser Arg Asn Arg Ser Phe Ser Lys Ala Ser Gly Pro Thr
                                    200
    209 Gln Tyr Ile Phe His Glu Glu Asp Met Asn Phe Val Asp Ala Pro Thr
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210
                                  215
                                                      220
     210
     211 Ile Ser Arg Val Phe Asp Glu Lys Thr Met Tyr Arg Asn Phe Ser Ser
                                                  235
     212 225
                             230
     213 Pro Arg Gly Met Cys Leu Ile Ile Asn Asn Glu His Phe Glu Gln Met
     214
                         245
                                              250
     215 Pro Thr Arg Asn Gly Thr Lys Ala Asp Lys Asp Asn Leu Thr Asn Leu
                     260
                                          265
                                                               270
     217 Phe Arg Cys Met Gly Tyr Thr Val Ile Cys Lys Asp Asn Leu Thr Gly
                 275
                                      280
     219 Arg Gly Met Leu Leu Thr Ile Arg Asp Phe Ala Lys His Glu Ser His
                                  295
             290
     221 Gly Asp Ser Ala Ile Leu Val Ile Leu Ser His Gly Glu Glu Asn Val
     222 305
     223 Ile Ile Gly Val Asp Asp Ile Pro Ile Ser Thr His Glu Ile Tyr Asp
                         325
                                              330
                                                                   335
     225 Leu Leu Asn Ala Ala Asn Ala Pro Arg Leu Ala Asn Lys Pro Lys Ile
                     340
                                       / 345
W--> 227 Val Phe Val Gln Ala Cys Arg Xaa Glu Arg Arg Asp Asn Gly Phe Pro
                                      360
     228
                 355
     229 Val Leu Asp Ser Val Asp Gly Val Pro Ala Phe Leu Arg Arg Gly Trp
     230
             370
                                  375
     231 Asp Asn Arg Asp Gly Pro Leu Phe Asn Phe Leu Gly Cys Val Arg Pro
     232 385
                             390
                                                  395
                  · /
  -> 233 Gln Val Xaa Gln Val Trp Arg Lys Lys Pro Ser Xaa Ala Asp Ile Leu
     234
                         405
    /235 Ile Arg Tyr Ala Thr Thr Ala Gln Tyr Val Ser Xaa Arg Asn Ser Ala
     236
                    420
                                          425
     237 Arg Gly Ser Trp Phe Ile Gln Ala Val Cys Glu Val Phe Ser Thr His
     238
                 435
W--> 239 Xaa Lys Asp Met Asp Val Val Glu Leu Leu Thr Glu Val Asn Lys Lys
     240
             450
                                  455
                                                      460
W--> 241 Val Xaa Cys Gly Phe Gln Thr Ser Gln Gly Ser Asn Ile Leu Lys Gln
                                                  475
                             470/
W--> 243 Met Pro Xaa Met Thr Xaa Arg Leu Leu Lys Lys Phe Tyr Phe Trp Pro
                         485
                                              490
     244
     245 Glu Ala Arg Asn Ser Ala Val
     246
                     500
     249 <210> SEQ ID NO: 3
     250 <211> LENGTH: 1373
     251 <212> TYPE: DNA
     252 <213> ORGANISM: Homo sapiens
     254 <220> FEATURE:
     255 <221> NAME/KEY: CDS
     256 <222> LOCATION: (18)...(1229)
     258 <400> SEQUENCE: 3
                                                                             50
     259 aaaaqqaqaq aaaaqcc atg gcc gac aag gtc ctg aag gag aag aga aag
                            Met Ala Asp Lys Val Leu Lys Glu Lys Arg Lys
     260
     261
                             1
                                                                             98
     263 ctg ttt atc cgt tcc atg ggt gaa ggt aca ata aat ggc tta ctg gat
```

Use of n and/or Kaa has been detected in the Sequence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/888,243

TIME: 13:33:10

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Input Set: A:\01997.211003.SEQLIST.TXT
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```
L:185 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:191 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:227 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:239 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:241 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:380 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:386 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:412 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:418 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:420 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:426 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
L:453 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:457 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:465 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:481 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
L:532 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:556 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:558 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:673 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:756 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:774 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16
L:806 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:824 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:865 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:869 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
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L:875 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:877 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:879 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 L:881 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
\dot{L}:900 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:902 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:904 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:906 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L\!:\!908 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:910 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:912 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:955 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L\!:\!990 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:992 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:996 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:998 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:1000 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
```

VERIFICATION SUMMARY

DATE: 02/07/2002

PATENT APPLICATION: US/09/888,243

TIME: 13:33:10

Input Set : A:\01997.211003.SEQLIST.TXT
Output Set: N:\CRF3\02072002\1888243.raw

L:1004 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 L:1023 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24